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S12 4	22	embedded with distribution with attributes	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	ON	2007/09/06 14:45
S12 7	0	717/104,106,168	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	ON	2007/09/11 17:08
S12 6	741	second with data adj model	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	ON	2007/09/11 17:08
S12 5	1444946	second with data model	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	ON	2007/09/11 17:08
S12 9	9570	((717/104,106,168) or (707/10)). CCLS.	US-PGPUB; USPAT; USOCR	OR	OFF	2007/09/11 17:09
S12 8	2126	(717/104,106,168).CCLS.	US-PGPUB; USPAT; USOCR	OR	OFF	2007/09/11 17:09
S13 2	22	embedded with distribution with attributes	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	ON	2007/09/12 16:38
S13 1	741	second with data adj model	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	ON	2007/09/12 16:38
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S13 6		S131 and S132	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	ON	2007/09/12 16:39
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L3	9829	((717/104-106,168) or (707/10)). CCLS.	US-PGPUB; USPAT; USOCR	OR	OFF	2007/09/12 21:04
L4	0	1 and 3	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	ON	2007/09/12 21:04
L5	40	2 and 3	US-PGPUB; USPAT; USOCR; EPO; JPO; DERWENT; IBM_TDB	OR	ON	2007/09/12 21:04
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May 2007 The Journal of Machine Learning Research, Volume 8

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This paper is about classifying entities that are interlinked with entities for which the class is known. After surveying prior work, we present NetKit, a modular toolkit for classification in networked data, and a case-study of its application to networked data used in prior machine learning research. NetKit is based on a node-centric framework in which classifiers comprise a local classifier, a relational classifier, and a collective inference procedure. Various existing node-centric relati ...

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Wai-Ho Au, Keith C. C. Chan, Andrew K. C. Wong, Yang Wang

April 2005 IEEE/ACM Transactions on Computational Biology and Bioinformatics (TCBB), Volume 2 Issue 2

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This paper presents an attribute clustering method which is able to group genes based on their interdependence so as to mine meaningful patterns from the gene expression data. It can be used for gene grouping, selection, and classification. The partitioning of a relational table into attribute subgroups allows a small number of attributes within or across the groups to be selected for analysis. By clustering attributes, the search dimension of a data mining algorithm is reduced. The reduction of  $\dots$ 

Keywords: Data mining, attribute clustering, gene selection, gene expression classification, microarray analysis.

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